

Amendments to the Specification:

Please replace the paragraph beginning at page 18, line 9 with the following amended paragraph:

OATP-C genotype data for 3 SNPs Asn130Asp, Pro 155Thr and Val174Ala was utilised to determine the haplotypes. The package SNPHAP (Clayton, David SNPHAPv0.2 2002 ~~<http://www-gene.cimr.cam.ac.uk/clayton/software/snphap.txt>~~ world wide web-gene at cimr.cam.ac.uk/clayton/software/snphap.txt) was used for this analysis. The haplotypes were also predicted using the PHASE package (Stephens, M. 2001 American Journal of Human Genetics 68, 978-989) and were found to give the same predicted haplotypes. The haplotypes were found to be concordant with those reported previously (Tirona 2001). The -118 A>C promoter SNP, at the NF1 binding site, was in strong linkage disequilibrium with the Val174Ala coding variant ( $\Delta=0.3$ ).

Please replace the paragraph beginning on page 18, line 17 with the following rewritten paragraph:

**Table 1 Common haplotypes in the OATP-C gene**

| <b>* nomenclature</b> | <b>Amino acid variant(s) on allele</b>                  | <b>Haplotype frequency (n=79)</b> |
|-----------------------|---|-----------------------------------|
| *1a                   | <del>Asp130</del> <u>Asn130</u> , Pro155, Val174        | 57%                               |
| *1b                   | <del>Asn130</del> <u>Asp130</u> , Pro155, Val174        | 22%                               |
| *5                    | <del>Asp130</del> <u>Asn130</u> , Pro155, <b>Ala174</b> | 2%                                |
| *14                   | <del>Asn130</del> <u>Asp130</u> , Thr155, Val174        | 7.5%                              |
| *15                   | <del>Asn130</del> <u>Asp130</u> , Pro155, <b>Ala174</b> | 11.5%                             |